Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

- 1-38. (previously canceled)
- 39. (currently amended) A An isolated cell, which is either an E. coli LacZ or a yeast, that produces a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, the cell comprising heterologous genes encoding one or more sugar nucleotide regenerating enzyme and one or more glycosyltransferase.
- 40. (canceled)
- 41. (canceled)
- 42. (canceled)
- 43. (currently amended) The <u>isolated</u> cell of claim-42 39, wherein the <u>isolated cell is</u> E. coli is LacZ.
- 44. (canceled)
- 45. (currently amended) The <u>isolated</u> cell of claim-44 <u>39</u>, wherein the <u>isolated</u> eukaryotic cell is a-yeast.
- 46. (currently amended) The <u>isolated</u> cell of claim 39, wherein at least one of the heterologous genes is integrated into the genome of the cell.
- 47. (currently amended) The <u>isolated</u> cell of claim 39, wherein the heterologous genes are encoded within one or more plasmids.

48. (currently amended) The <u>isolated</u> cell of claim 47, wherein the heterologous genes are encoded within one plasmid.

- 49. (canceled)
- 50-51. (previously canceled)
- 52. (currently amended) The isolated cell of claim 39, wherein the one or more sugar nucleotide regenerating enzyme is selected from the group consisting of galactokinase, galactose-1-phosphate uridylyltransferase, glucose-1-phosphate uridylyltransferase, pyruvate kinase, nucleotide diphosphate kinase, polyphosphate kinase, acetate kinase, pyruvate oxidase, pyrophosphatase, phosphoglucomutase, N-acetylglucosamine permease, acetylglucosaminephosphate mutase, N-acetylglucosamine-1-phosphate uridyltransferase, Nacetylglucosamine kinase, pyrophosphorylase, uridine 5'-diphosphoglucouronic acid 6-dehydrogenase, N-acetylneuraminate lyase, sialic acid aldolase, cytosine 5'-monophosphate kinase, cytosine 5'-monophosphate-N-acetylneuraminic acid synthetase, α 1,3-mannosyltransferase, guanosine 5'-diphosphomannose:Dol-PP-N-acetylglucosamine β-mannosyltransferase, sucrose synthase synthetase, mannose-1-phosphate guanyltransferase, guanisine 5'-diphosphate-mannose pyrophosphorlyase, a phosphomannomutase, uridine 5'-diphosphate-galactose 4epimease, uridine 5'-diphosphate-glucose 4-epimerase, guanosine 5'-diphosphatemannose pyrophosphorylase, guanosine 5'-diphosphate-D-mannose 4.6deydratase, and guanosine 5'-diphosphate-L-fucose synthetase.
- 53. (currently amended) The <u>isolated</u> cell of claim 39 comprising genes encoding galactokinase, galactose-1-phosphate uridylyltransferase, and glucose-1-phosphate uridylyltransferase.
- 54. (currently amended) The <u>isolated</u> cell of claim 39 comprising a gene encoding nucleotide diphosphate kinase.
- 55. (currently amended) The <u>isolated</u> cell of claim 53 comprising a gene encoding polyphosphate kinase.

56. (currently amended) The <u>isolated</u> cell of claim 53 comprising a gene encoding pyruvate kinase.

- 57. (currently amended) The <u>isolated</u> cell of claim 53 comprising genes encoding pyruvate oxidase, nucleotide diphosphate kinase, and pyrophosphatase.
- 58. (currently amended) The <u>isolated</u> cell of claim 39 comprising a gene encoding sucrose synthetase.
- 59. (currently amended) The <u>isolated</u> cell of claim 58 further comprising a gene encoding uridine 5'-diphosphate-galactose 4-epimease or uridine 5'-diphosphate-glucose 4-epimerase.
- 60. (currently amended) The <u>isolated</u> cell of claim 58 further comprising a gene encoding glucosyltransferase.
- 61. (currently amended) The <u>isolated</u> cell of claim 58 further comprising genes encoding uridine 5'-diphosphoglucouronic acid 6-dehydrogenase and a glucuronyltransferase.
- 62. (currently amended) The <u>isolated</u> cell of claim 39, wherein the one or more glycosyltransferase(s) is selected from the group consisting of a galactosyltransferase, a glucosyltransferase, a N-acetylglucosaminyl transferase, an N-acetylgalactosaminyl transferase, a glucuronyltransferase, a sialyltransferase, a mannosyltransferase, and a fucosyltransferase.
- 63. (currently amended) The <u>isolated</u> cell of claim 62 wherein the galactosyltransferase is selected from the group consisting of α1,3 galactosyltransferase, β1,4 galactosyltransferase, and α1,4 galactosyltransferase.
- 64. (currently amended) The <u>isolated</u> cell of claim 62, wherein the glycosyltransferase is a glucosyltransferase.
- 65. (currently amended) The <u>isolated</u> cell of claim 62, wherein the glycosyltransferase is a N-acetylglucosaminyl transferase.

66. (currently amended) The <u>isolated</u> cell of claim 62, wherein the N-acetylgalactosaminyl transferase is uridine 5'-diphosphate-N-acetylgalactosamine:2'-fucosylgalactoside-α-3-N-acetylgalctosaminyl transferase.

- 67. (currently amended) The <u>isolated</u> cell of claim 62, wherein the glycosyltransferase is a glucuronyltransferase.
- 68. (currently amended) The <u>isolated</u> cell of claim 62, wherein the glycosyltransferase is a sialyltransferase.
- 69. (currently amended) The <u>isolated</u> cell of claim 62, wherein the glycosyltransferase is a mannosyltransferase.
- 70. (currently amended) The <u>isolated</u> cell of claim 62, wherein the fucosyltransferase is selected from the group consisting of $\alpha 1,3$ fucosyltransferase, $\alpha 1,2$ fucosyltransferase, and $\alpha 1,3/4$ fucosyltransferase.

71-76. (canceled)